

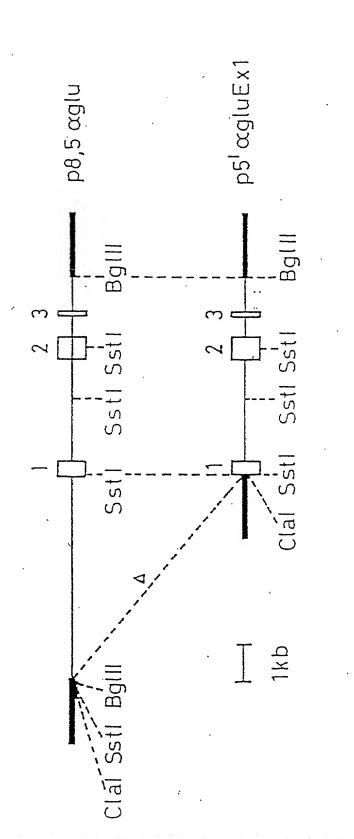
 $lpha_{f S_1}$ casein sequence, promoler or 3' untranslated region.

The boxes represent the exons in the α -glucosidase sequence, the thin line represents the intron sequences. The numbers above the boxes are the exon numbers

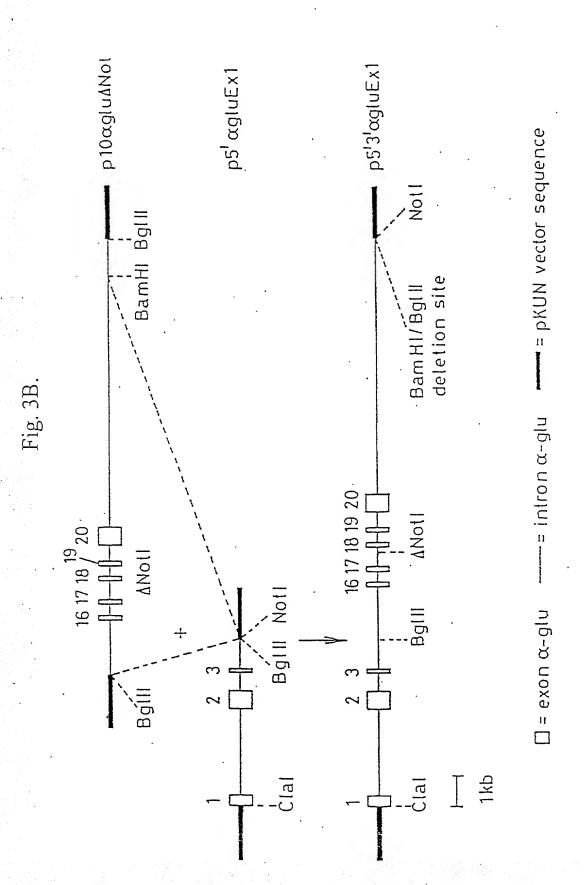
The numbers above the boxes are to polyadenylation signal.

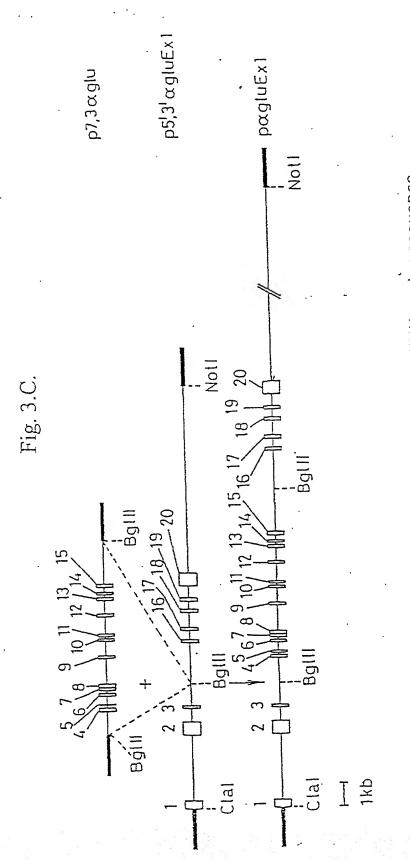
ATG = translation initiation sile, TAG = translation stop codon

Fig. 3A.

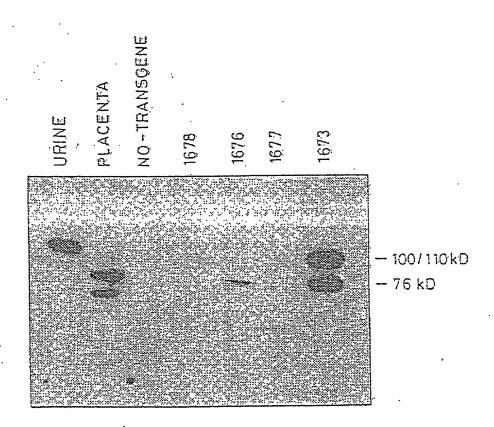


= pKUN vector sequence = intron \alpha-glu = exon ∞ -glu

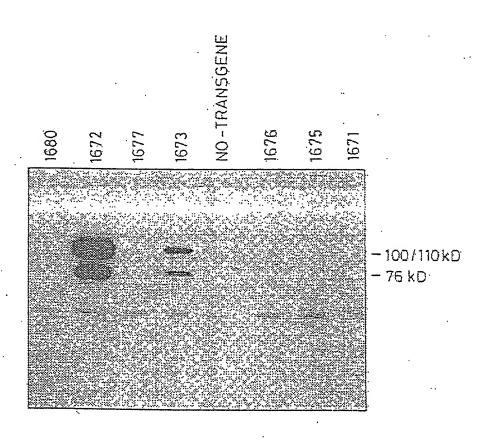


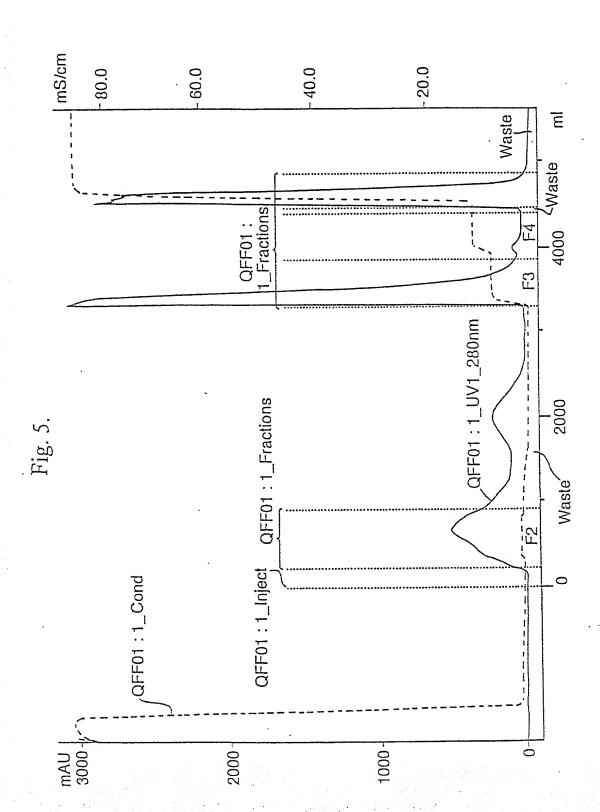


= pKUN vector sequence = intron a-glu □ = exon α-glu

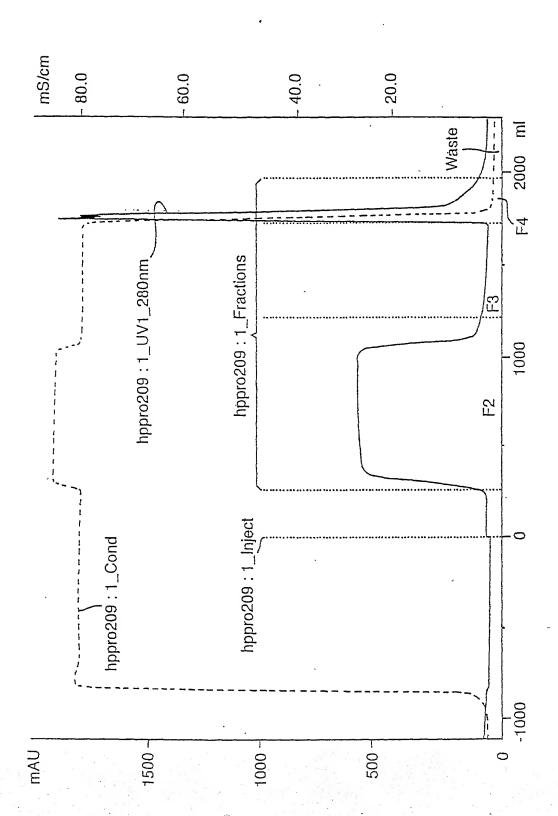












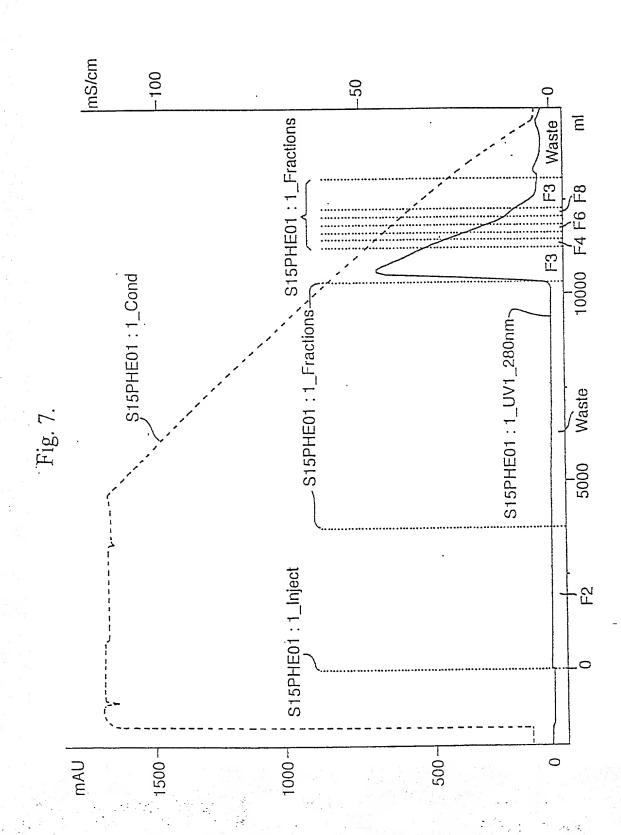


Fig. 8.

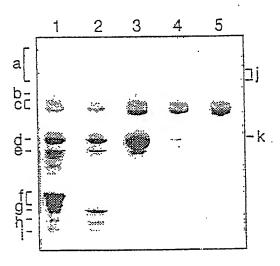
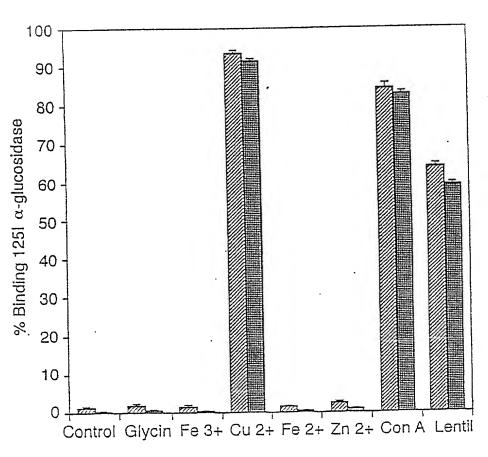


Fig. 9. -8.40 , O C Judanhanhari JA-GLU-13 14 12 5 8 9 11 6 7 9 10 Time [min]

Fig. 10.



PBS, 0.1% Tween-20, 0.5 M NaCl wash

PBS, 0.02% Tween-20 wash

Fig. 11. A.

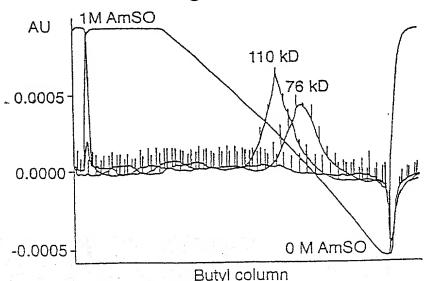


Fig. 11. B.

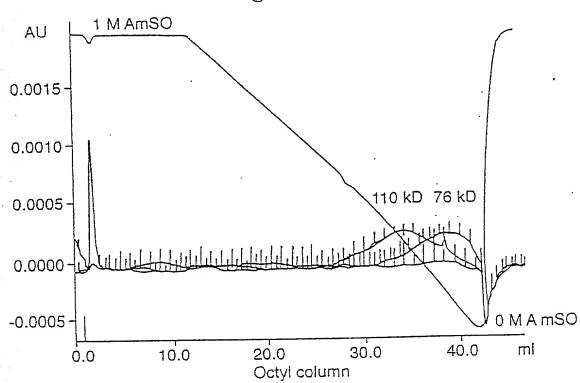


Fig. 11. C.

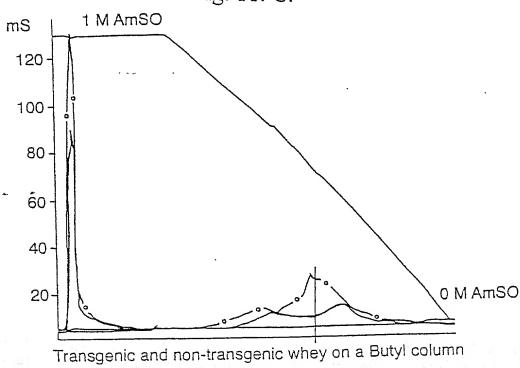
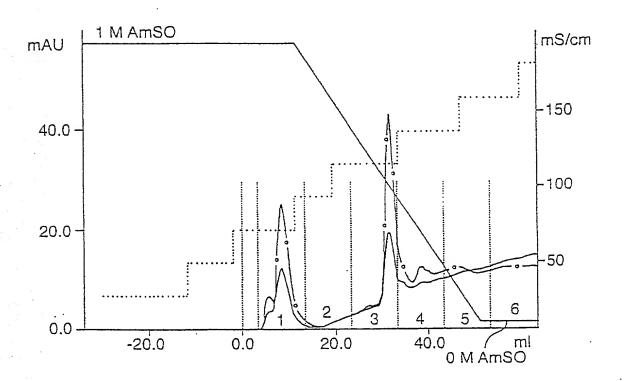
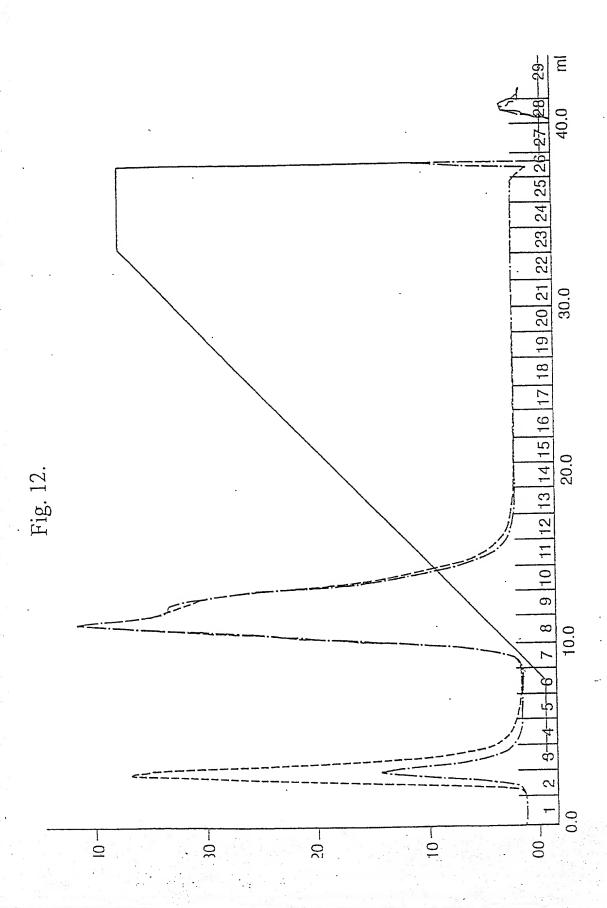
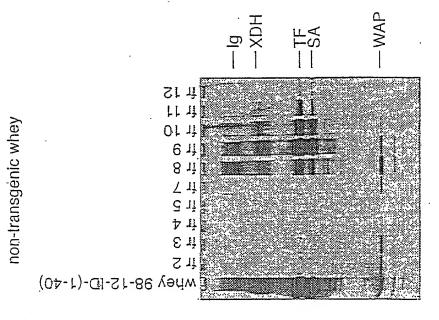


Fig. 11. D.







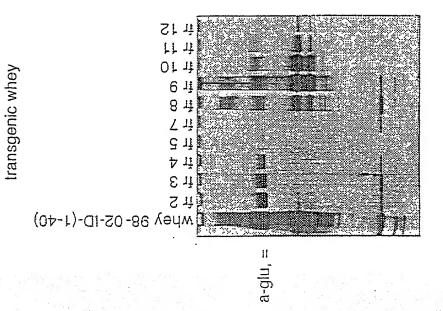
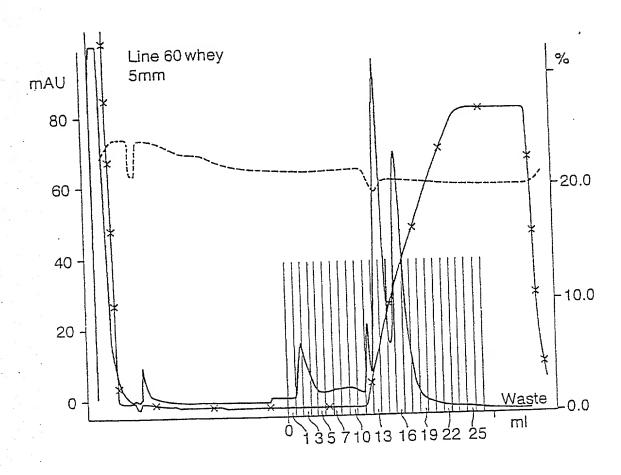


Fig. 14.



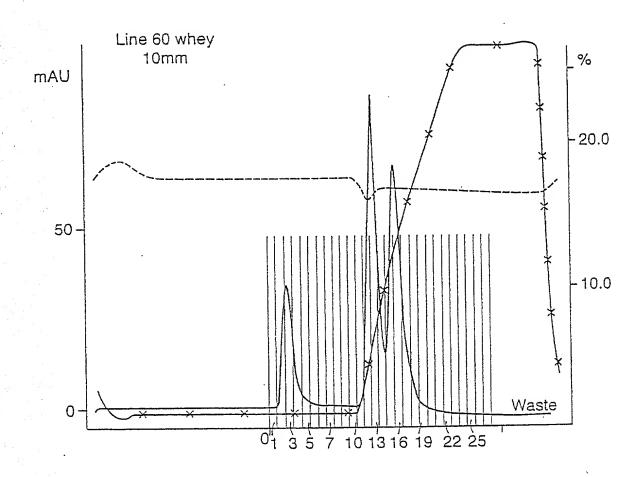
-12099801:1_UV1_280nm

_12099801:1_pH

×12099801:1_Cond%

12099801:1_Fractions

Fig. 15.



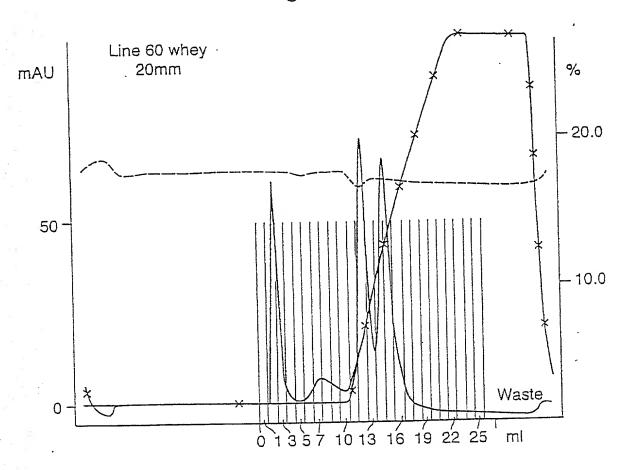
_____ 12099802:11_UV1_280nm

---- 12099802:11_pH

-×--×- 12099802:11_Cond%

12099802:11_Fractions

Fig. 16.



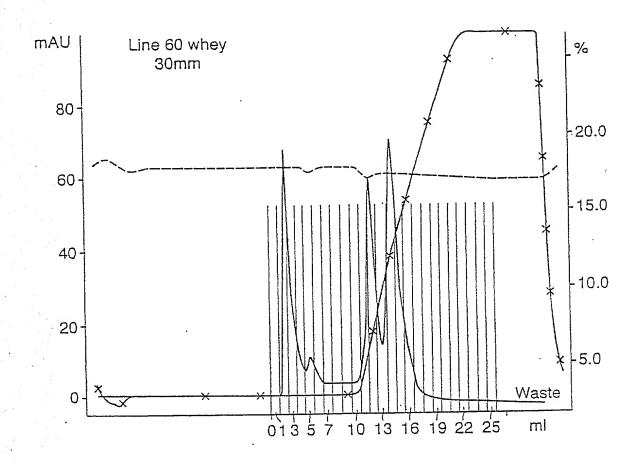
12099803:12_UV1_280nm

12099803:12_pH

12099803:12_Cond%

12099803:12_Fractions

Fig. 17.



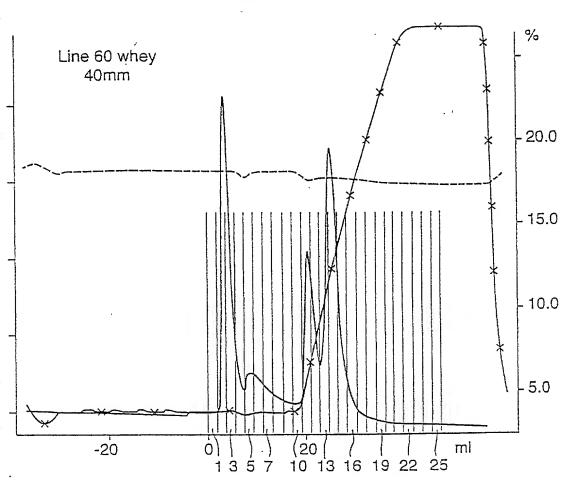
12099804:13_UV1_280nm

12099804:13_pH

12099804:13_Cond%

12099804:13_Fractions

Fig. 18.



121099805:1_UV1_280nm

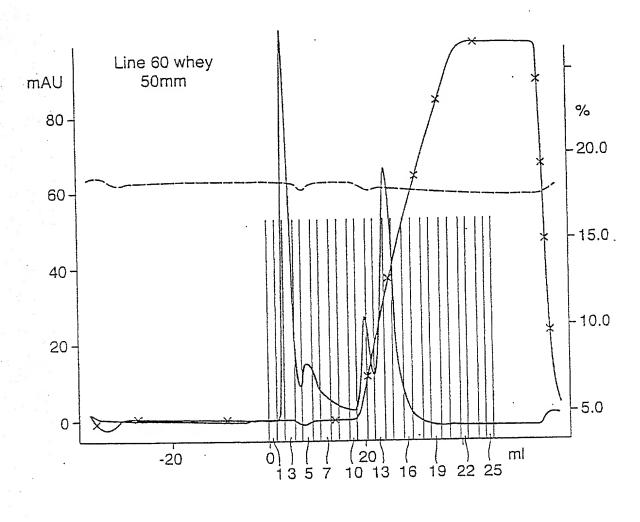
---- 121099805:1_pH

** * * 121099805:1_Cond%

121099805:1_Fractions

TOTAL TATE

Fig. 19.



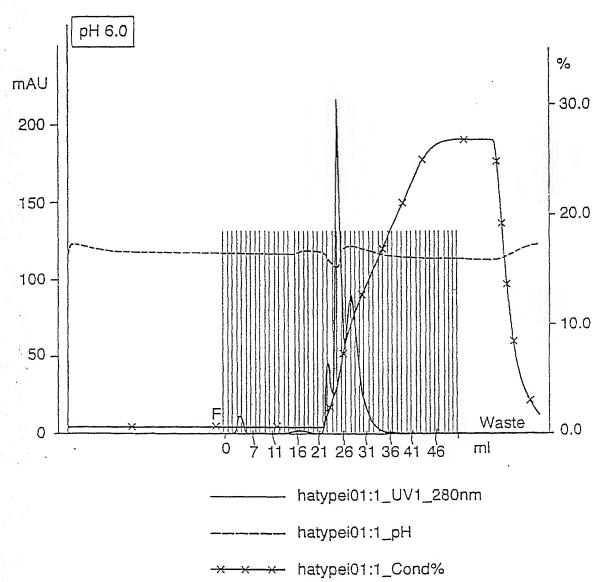
121099806:1_UV1_280nm

121099806:1_pH

× 121099806:1_Cond%

121099806:1_Fractions

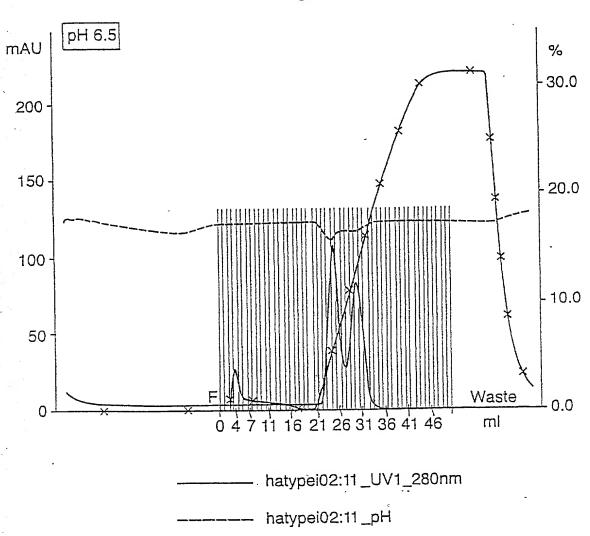
Fig. 20.



hatypei01:1_Fractions

with the train with the train with the train with

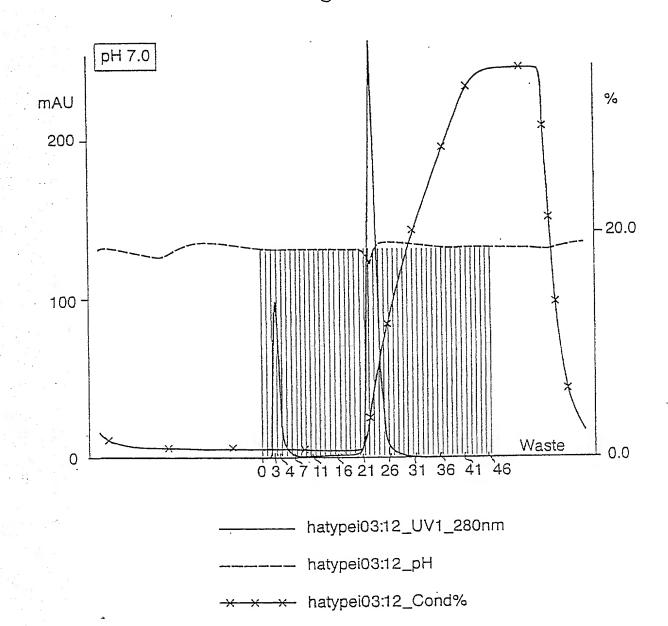
Fig. 21.



x x x hatypei02:11_Cond%

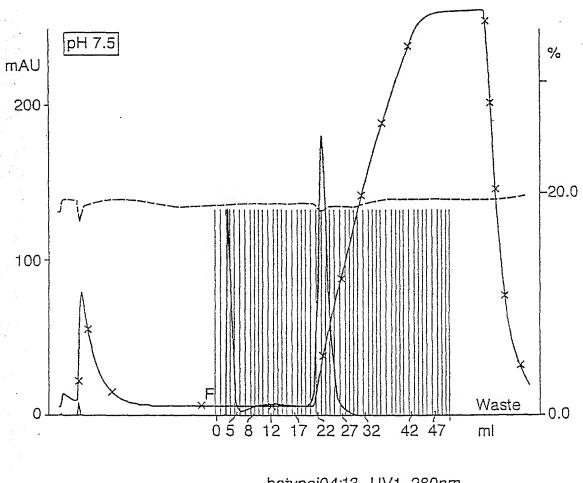
hatypei02:11 _ Fractions

Fig. 22.



hatypei03:12_Fractions

Fig. 23.



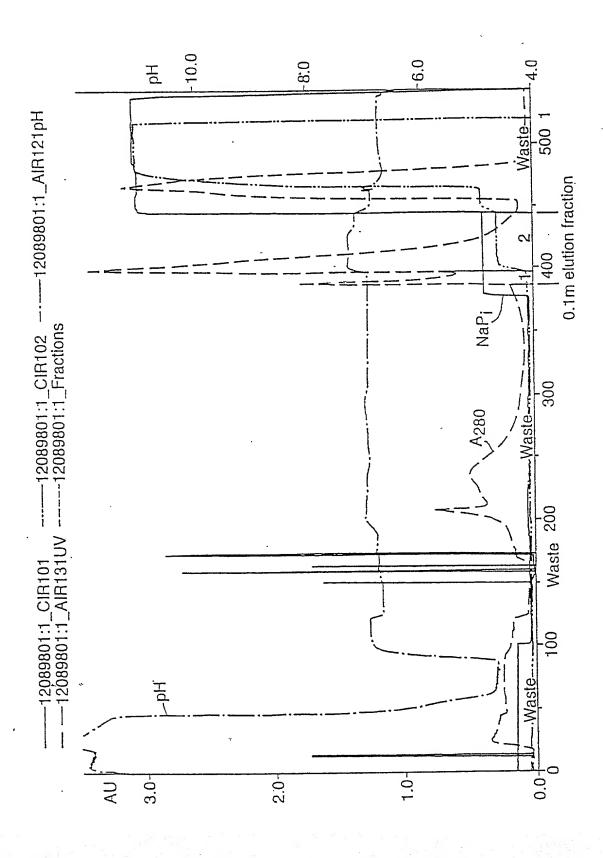
hatypei04:13_UV1_280nm

hatypei04:13_pH

hatypei04:13_Cond%

hatypei04:13 _ Fractions







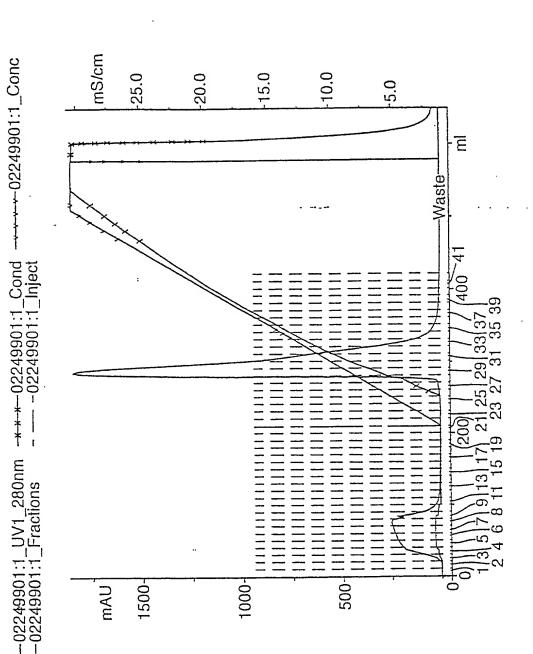


Fig. 26.

XK16/15 80°C cHT type I 10mM Napi pH 6.5 ; QFF eluate Run 02249901/02259901/02269901

